

Figure 1A

>SEQ\_ID\_NO:1\_SGP057

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CCCCACGGCCGGATGAGAAGTCTCTCCCTGCAGCCCTCAAGGCCACGAGGATCAGTGCTCGGA  
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GAGACTAGGAGTCGAGCAGCTGCGGCTCAGCACAGTAGACATGACTGGGATCCCCACCTTGGACA  
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Figure 1B

>SEQ\_ID\_NO:4\_SGP045

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Figure 2

>SEQ\_ID\_NO:6\_SGP057

RWFHPNISRVEAEKLFSLRGQRGDFLARPSSESPGGFTLSVREALGEGSLSQGLSRTLSSPVRALGTGRA  
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SDPGGFPLSALTQGWDEAQGSGRQPQVTHIMTHSQVGGGGELGRPLGRAGGLGQAPHRHPHGRMRSL  
SLQPLKATRISARSLEGCQVESRATDASGKARQGFVVEFKMLQQQECRFLYPRKEGQSVENKPKNIHY  
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FTRLEFPRLKGGDLGQKVLFRDHHMSGWSYKRVEKSDLKYPLIHGQGRQARLLGTLAVSRGLGDHQ  
LRVLDTNILKPFLLSVPQVTVLDVDQLELQEDDVVVVMTDGLWDVLSNEQVAWLVRSLPGNQEDP  
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GSLDNMTCILVCFPGAPRPSEEAIRRELALDAALGCRIELCASAQKPPSLNTVFRTLASEDIPDLPPGGG  
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SRDNI